SEQUENCE LISTING

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<110>	Statens Serum Intitut Anderson, Peter
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<140>	09/804,980

<160> 257

<141> 2001-03-12

<170> PatentIn version 3.0

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Ile Thr Tyr Gln Ala Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Asp 50 55 60

Leu Val Arg Ala Tyr His Ala Met Ser Ser Thr His Glu Ala Asn Thr



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Gly Leu Asp Val Ser Asp Arg Ile Arg Val Val Met Ser Val Pro Ala 50 55 60

Glu Arg Glu Asp Trp Ala Arg Thr His Arg Asp Leu Ile Ala Gly Glu 65 70 75 80

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Glu Gly Leu Pro Pro Gly Ser Ala Leu Leu Val Val Leu Arg Gly Pro 50 55 60

Asn Ala Gly Ser Arg Pro Leu Leu Asp Gln Ala Ile Thr Ser Ala Gly 65 70 75 80

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Glu	Asp	Tyr 115	Gly	Val	Thr	Ile	Ala 120	Asp	Gly	Pro	Met	Ala 125	Gly	Leu	Leu	
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Ala Ser Gly Leu Gly Asp Val Gly Glu Ala Phe Val Asp Ser Leu Thr 50 55 60

Ser Gln Val Gly Gly Arg Ser Ile Gly Val Tyr Ala Val Asn Tyr Pro 75 80

Ala Ser Asp Asp Tyr Arg Ala Ser Ala Ser Asn Gly Ser Asp Asp Ala 85 90 95

Ser Ala His Ile Gln Arg Thr Val Ala Ser Cys Pro Asn Thr Arg Ile 100 105 110

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<211> 217

<212> PRT

<213> Mycobacterium tuberculosis

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Pro Gly Ile Gly Thr Val Gly Asn Ala Phe Val Ser Ala Leu Arg Ser 50 55 60

Lys Val Asn Lys Asn Val Gly Val Tyr Ala Val Lys Tyr Pro Ala Asp 65 70 75 80

Asn Gln Ile Asp Val Gly Ala Asn Asp Met Ser Ala His Ile Gln Ser 85 90 95

Met Ala Asn Ser Cys Pro Asn Thr Arg Leu Val Pro Gly Gly Tyr Ser 100 105 110

Leu Gly Ala Ala Val Thr Asp Val Val Leu Ala Val Pro Thr Gln Met

115 120 125

Trp Gly Phe Thr Asn Pro Leu Pro Pro Gly Ser Asp Glu His Ile Ala 130 135 140

Ala Val Ala Leu Phe Gly Asn Gly Ser Gln Trp Val Gly Pro Ile Thr 145 150 155 160

Asn Phe Ser Pro Ala Tyr Asn Asp Arg Thr Ile Glu Leu Cys His Gly
165 170 175

Asp Asp Pro Val Cys His Pro Ala Asp Pro Asn Thr Trp Glu Ala Asn 180 185 190

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Glu Gly I	le Arg Sei	Ala Ser 135		sn Pro		Leu Thr 140	Leu	Pro	Glu				
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Arg Leu A	la Gly Vai 16!		Pro T	yr Ser 170	Val 1	Leu Leu	Ser	Ala 175	Asp				
Val Tyr T	nr Lys Val	l Ser Glu		er Asp 85	His (Gly Tyr	Pro 190	Ile	Arg				
Glu His L	eu Asn Arg	g Leu Val	Asp G:	ly Asp	Ile :	Ile Trp 205	Ala	Pro	Ala				

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210 215 220

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Ala Gly Gly Ala Tyr Ser Met Tyr Thr Asn Trp Glu Gln Asp Gly Ser 100 105 110

Lys Gln Trp Asp Thr Phe Leu Ser Ala Glu Leu Pro Asp Trp Leu Ala 115 120 125

Ala Asn Arg Gly Leu Ala Pro Gly Gly His Ala Ala Val Gly Ala Ala 130 135 140

Gln Gly Gly Tyr Gly Ala Met Ala Leu Ala Ala Phe His Pro Asp Arg 145 150 155 160

Phe Gly Phe Ala Gly Ser Met Ser Gly Phe Leu Tyr Pro Ser Asn Thr 165 170 175

Thr Thr Asn Gly Ala Ile Ala Ala Gly Met Gln Gln Phe Gly Gly Val 180 185 190

Asp Thr Asn Gly Met Trp Gly Ala Pro Gln Leu Gly Arg Trp Lys Trp 195 200 205

His Asp Pro Trp Val His Ala Ser Leu Leu Ala Gln Asn Asn Thr Arg 210 215 220

Val Trp Val Trp Ser Pro Thr Asn Pro Gly Ala Ser Asp Pro Ala Ala 225 230 235 240

Met Ile Gly Gln Thr Ala Glu Ala Met Gly Asn Ser Arg Met Phe Tyr
245 250 255

Asn Gln Tyr Arg Ser Val Gly Gly His Asn Gly His Phe Asp Phe Pro 260 265 270

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<212> DNA

<213> Mycobacterium tuberculosis

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Leu Leu Thr Pro Ala Leu Ala Pro Pro Ala Ser Ala Gly Cys Pro 35 40 45

Asp Ala Glu Val Val Phe Ala Arg Gly Thr Gly Glu Pro Pro Gly Leu 50 55 60

Gly Arg Val Gly Gln Ala Phe Val Ser Ser Leu Arg Gln Gln Thr Asn 65 70 75 80

Lys Ser Ile Gly Thr Tyr Gly Val Asn Tyr Pro Ala Asn Gly Asp Phe 85 90 95

Leu Ala Ala Asp Gly Ala Asn Asp Ala Ser Asp His Ile Gln Gln
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Met Ala Ser Ala Cys Arg Ala Thr Arg Leu Val Leu Gly Gly Tyr Ser 115 120 125

Gln Gly Ala Ala Val Ile Asp Ile Val Thr Ala Ala Pro Leu Pro Gly
130 135 140

Leu Gly Phe Thr Gln Pro Leu Pro Pro Ala Ala Asp Asp His Ile Ala 145 150 155 160

Ala Ile Ala Leu Phe Gly Asn Pro Ser Gly Arg Ala Gly Gly Leu Met 165 170 175

Ser Ala Leu Thr Pro Gln Phe Gly Ser Lys Thr Ile Asn Leu Cys Asn 180 185 190

Asn Gly Asp Pro Ile Cys Ser Asp Gly Asn Arg Trp Arg Ala His Leu

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Arg Ile 225

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<211> 700

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Tyr Ser Gln Ala Val Glu Glu Arg Asn His Ala Met Met Leu Val Gln 50 55 60

His Leu Leu Asp Arg Asp Leu Arg Val Glu Ile Pro Gly Val Asp Thr 65 70 75 80

Val Arg Asn Gln Phe Asp Arg Pro Arg Glu Ala Leu Ala Leu 85 90 95

Asp Gln Glu Arg Thr Val Thr Asp Gln Val Gly Arg Leu Thr Ala Val
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Ala Arg Asp Glu Gly Asp Phe Leu Gly Glu Gln Phe Met Gln Trp Phe
115 120 125

Leu Gln Glu Gln Ile Glu Glu Val Ala Leu Met Ala Thr Leu Val Arg 130 135 140

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Gly Val Tyr Pro Val Asn Tyr Ala Ala Ser Arg Leu Gln Leu His Gly 85 90 95
Gly Asp Gly Ala Asn Asp Ala Ile Ser His Ile Lys Ser Met Ala Ser 100 105 110
Ser Cys Pro Asn Thr Lys Leu Val Leu Gly Gly Tyr Ser Gln Gly Ala 115 120 125
Thr Val Ile Asp Ile Val Ala Gly Val Pro Leu Gly Ser Ile Ser Phe 130 135 140
Gly Ser Pro Leu Pro Ala Ala Tyr Ala Asp Asn Val Ala Ala Val Ala 145 150 155 160
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Pro Ile Cys His Val Gly Pro Gly Asn Glu Phe Ser Gly His Ile Asp 195 200 205
Gly Tyr Ile Pro Thr Tyr Thr Gln Ala Ala Ser Phe Val Val Gln 210 215 220
Arg Leu Arg Ala Gly Ser Val Pro His Leu Pro Gly Ser Val Pro Gln

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Tyr Pro Gly Gly Val Val Met Ala Gly Asp Arg Arg Ser Thr Gln Gly 70

Asn Met Ile Ser Gly Arg Asp Val Arg Lys Val Tyr İle Thr Asp Asp

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Ala Arg Leu Tyr Ala Val Glu Leu Glu His Tyr Glu Lys Leu Glu Gly 120

Val Pro Leu Thr Phe Ala Gly Lys Ile Asn Arg Leu Ala Ile Met Val 135

Arg Gly Asn Leu Ala Ala Ala Met Gln Gly Leu Leu Ala Leu Pro Leu 150

Leu Ala Gly Tyr Asp Ile His Ala Ser Asp Pro Gln Ser Ala Gly Arg 170

Ile Val Ser Phe Asp Ala Ala Gly Gly Trp Asn Ile Glu Glu Gly 180 185 190

Tyr Gln Ala Val Gly Ser Gly Ser Leu Phe Ala Lys Ser Ser Met Lys 200

Lys Leu Tyr Ser Gln Val Thr Asp Gly Asp Ser Gly Leu Arg Val Ala 215

Val Glu Ala Leu Tyr Asp Ala Ala Asp Asp Asp Ser Ala Thr Gly Gly 225 230

Pro Asp Leu Val Arg Gly Ile Phe Pro Thr Ala Val Ile Ile Asp Ala 250

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tuberculosis

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gcgcgcgttc	cggcgcatca	ccggctccgc	cctgcaagcg	ttgctggtag	accaggaaag	780
cccgcagtct	gacggcgaat	cgtcgggctg	agtccgaaag	tccgacgcgt	gtctgggacc	840
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<210> 60

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Leu Ala Tyr Ala Gly Gly Val Leu Phe Val Ala Glu Asn Pro Ser Arg 40

Ser Leu Gln Lys Ile Ser Glu Leu Tyr Asp Arg Val Gly Phe Ala Ala 50

Ala Gly Lys Phe Asn Glu Phe Asp Asn Leu Arg Arg Gly Gly Ile Gln 70 75

<211> 248

<212> PRT

<213> Mycobacterium tuberculosis

<400> 60

Phe Ala Asp Thr Arg Gly Tyr Ala Tyr Asp Arg Arg Asp Val Thr Gly 85 90 95	,				
Arg Gln Leu Ala Asn Val Tyr Ala Gln Thr Leu Gly Thr Ile Phe Thr 100 105 110	;				
Glu Gln Ala Lys Pro Tyr Glu Val Glu Leu Cys Val Ala Glu Val Ala 115 120 125	ı				
His Tyr Gly Glu Thr Lys Arg Pro Glu Leu Tyr Arg Ile Thr Tyr Asp 130 135 140)				
Gly Ser Ile Ala Asp Glu Pro His Phe Val Val Met Gly Gly Thr Thr 145 150 155 160					
Glu Pro Ile Ala Asn Ala Leu Lys Glu Ser Tyr Ala Glu Asn Ala Ser 165 170 175	:				
Leu Thr Asp Ala Leu Arg Ile Ala Val Ala Ala Leu Arg Ala Gly Ser 180 185 190	=				
Ala Asp Thr Ser Gly Gly Asp Gln Pro Thr Leu Gly Val Ala Ser Leu 195 200 205	1				
Glu Val Ala Val Leu Asp Ala Asn Arg Pro Arg Arg Ala Phe Arg Arg 210 215 220	9				
Ile Thr Gly Ser Ala Leu Gln Ala Leu Leu Val Asp Gln Glu Ser Pro 225 230 235 240					
Gln Ser Asp Gly Glu Ser Ser Gly 245					
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totoggagoc ggtocoggog ggtatgtogo ggcgattogo gccgcacago toggoctg					
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attraggate agggggggg tgaggttggg staggggate gggtatggg aggcgaa					
atttggcatc agcggcgagg tgaccttcga ctacggcatc gcctatgacc gcagccga					
gatecaeggg taeggeacat ttgeegaege caacaegttg ttggttgate teaaegae					
January Control of the Control of th	-33 -300				

600 ggttcccggc acctcactgt cggccaacgt agtcacctac gaggaacaga tcctgtcccg agagetgeeg aaategatea ttattgeegg agetggtgee attggeatgg agtteggeta 660 cgtgctgaag aactacggcg ttgacgtgac catcgtggaa ttccttccgc gggcgctgcc 720 caacgaggac gccgatgtgt ccaaggagat cgagaagcag ttcaaaaagc tgggtgtcac 780 gatectgace gecaegaagg tegagteeat egeegatgge gggtegeagg teacegtgae 840 900 cgtcaccaag gacggcgtgg cgcaagagct taaggcggaa aaggtgttgc aggccatcgg atttgcgccc aacgtcgaag ggtacgggct ggacaaggca ggcgtcgcgc tgaccgaccg 960 1020 caaggctatc ggtgtcgacg actacatgcg taccaacgtg ggccacatct acgctatcgg 1080 cgatgtcaat ggattactgc agctggcgca cgtcgccgag gcacaaggcg tggtagccgc cgaaaccatt geeggtgeag agaetttgae getgggegae categgatgt tgeegegege 1140 gacgttctgt cagccaaacg ttgccagctt cgggctcacc gagcagcaag cccgcaacga 1200 aggttacgac gtggtggtgg ccaagttccc gttcacggcc aacgccaagg cgcacggcgt 1260 gggtgacccc agtgggttcg tcaagctggt ggccgacgcc aagcacggcg agctactggg 1320 tgggcacctg gtcggccacg acgtggccga gctgctgccg gagctcacgc tggcgcagag 1380 gtgggacctg accgccagcg agctggctcg caacgtccac acccacccaa cgatgtctga 1440 ggcgctgcag gagtgcttcc acggcctggt tggccacatg atcaatttct gagcggctca 1500 tgacgaggcg cgcgagcact gacaccccc agatcatcat gggtgccatc ggtggtgtgg 1560

Met Thr His Tyr Asp Val Val Leu Gly Ala Gly Pro Gly Gly Tyr 1 5 10 15

Val Ala Ala Ile Arg Ala Ala Gln Leu Gly Leu Ser Thr Ala Ile Val 20 25 30

Glu Pro Lys Tyr Trp Gly Gly Val Cys Leu Asn Val Gly Cys Ile Pro 35 40 45

Ser Lys Ala Leu Leu Arg Asn Ala Glu Leu Val His Ile Phe Thr Lys 50 55 60

Asp Ala Lys Ala Phe Gly Ile Ser Gly Glu Val Thr Phe Asp Tyr Gly 65 70 75 80

<210> 62

<211> 464

<212> PRT

<213> Mycobacterium tuberculosis

<400> 62

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- Val His Phe Leu Met Lys Lys Asn Lys Ile Thr Glu Ile His Gly Tyr
 100 105 110
- Gly Thr Phe Ala Asp Ala Asn Thr Leu Leu Val Asp Leu Asn Asp Gly
 115 120 125
- Gly Thr Glu Ser Val Thr Phe Asp Asn Ala Ile Ile Ala Thr Gly Ser 130 135 140
- Ser Thr Arg Leu Val Pro Gly Thr Ser Leu Ser Ala Asn Val Val Thr 145 150 155 160
- Tyr Glu Glu Gln Ile Leu Ser Arg Glu Leu Pro Lys Ser Ile Ile Ile 165 170 175
- Ala Gly Ala Gly Ala Ile Gly Met Glu Phe Gly Tyr Val Leu Lys Asn 180 185 190
- Tyr Gly Val Asp Val Thr Ile Val Glu Phe Leu Pro Arg Ala Leu Pro 195 200 205
- Asn Glu Asp Ala Asp Val Ser Lys Glu Ile Glu Lys Gln Phe Lys Lys 210 215 220
- Leu Gly Val Thr Ile Leu Thr Ala Thr Lys Val Glu Ser Ile Ala Asp 225 230 235 240
- Gly Gly Ser Gln Val Thr Val Thr Lys Asp Gly Val Ala Gln
 245 250 255
- Glu Leu Lys Ala Glu Lys Val Leu Gln Ala Ile Gly Phe Ala Pro Asn 260 265 270
- Val Glu Gly Tyr Gly Leu Asp Lys Ala Gly Val Ala Leu Thr Asp Arg 275 280 285
- Lys Ala Ile Gly Val Asp Asp Tyr Met Arg Thr Asn Val Gly His Ile 290 295 300
- Tyr Ala Ile Gly Asp Val Asn Gly Leu Leu Gln Leu Ala His Val Ala 305 310 315 320
- Glu Ala Gln Gly Val Val Ala Ala Glu Thr Ile Ala Gly Ala Glu Thr 325 330 335
- Leu Thr Leu Gly Asp His Arg Met Leu Pro Arg Ala Thr Phe Cys Gln 340 345 350
- Pro Asn Val Ala Ser Phe Gly Leu Thr Glu Gln Gln Ala Arg Asn Glu
 355 360 365
- Gly Tyr Asp Val Val Val Ala Lys Phe Pro Phe Thr Ala Asn Ala Lys 370 375 380

Ala His Gly Val Gly Asp Pro Ser Gly Phe Val 385 390 395	Lys Leu Val Ala Asp 400	
Ala Lys His Gly Glu Leu Leu Gly Gly His Leu 405 410	Val Gly His Asp Val 415	
Ala Glu Leu Leu Pro Glu Leu Thr Leu Ala Gln 420 425	Arg Trp Asp Leu Thr 430	
Ala Ser Glu Leu Ala Arg Asn Val His Thr His 435 440	Pro Thr Met Ser Glu 445	
Ala Leu Gln Glu Cys Phe His Gly Leu Val Gly 450 455	His Met Ile Asn Phe 460	
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cgaactgctg gacgcgttca aggaaatgac cctgttggag	ctctccgact tcgtcaagaa	180
gttcgaggag accttcgagg tcaccgccgc cgctccagtc	gccgtcgccg ccgccggtgc	240
cgcccggcc ggtgccgccg tcgaggctgc cgaggagcag	tccgagttcg acgtgatcct	300
tgaggccgcc ggcgacaaga agatcggcgt catcaaggtg	gtccgggaga tcgtttccgg	360
cctgggcctc aaggaggcca aggacctggt cgacggcgcg	cccaagccgc tgctggagaa	420
ggtcgccaag gaggccgccg acgaggccaa ggccaagctg	gaggccgccg gcgccaccgt	480
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<210> 64 <211> 130 <212> PRT <213> Mycobacterium tuberculosis		
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Thr Leu Leu Glu Leu Ser Asp Phe Val Lys Lys	Phe Glu Glu Thr Phe	

Glu Val Thr Ala Ala Ala Pro Val Ala Val Ala Ala Ala Gly Ala Ala

35 40 45

Pro Ala Gly Ala Ala Val Glu Ala Ala Glu Glu Gln Ser Glu Phe Asp 55 Val Ile Leu Glu Ala Ala Gly Asp Lys Lys Ile Gly Val Ile Lys Val Val Arg Glu Ile Val Ser Gly Leu Gly Leu Lys Glu Ala Lys Asp Leu 90 Val Asp Gly Ala Pro Lys Pro Leu Leu Glu Lys Val Ala Lys Glu Ala 100 Ala Asp Glu Ala Lys Ala Lys Leu Glu Ala Ala Gly Ala Thr Val Thr 120 Val Lys 130 <210> 65 <211> 900 <212> DNA <213> Mycobacterium tuberculosis <400> 65 tgaacgccat cgggtccaac gaacgcagcg ctacctgatc accaccgggt ctgttagggc 60 tettececag gtegtacagt egggecatgg ceattgaggt tteggtgttg egggttttea 120 ccgattcaga cgggaatttc ggtaatccgc tgggggtgat caacgccagc aaggtcgaac 180 accgcgacag gcagcagctg gcagcccaat cgggctacag cgaaaccata ttcqtcqatc 240 ttcccagccc cggctcaacc accgcacacg ccaccatcca tactccccqc accqaaattc 300 cgttcgccgg acacccgacc gtgggagcgt cctggtggct gcgcgagagg gggacgccaa 360 ttaacacgct gcaggtgccg gccggcatcg tccaggtgag ctaccacggt gatctcaccg 420 ccatcagcgc ccgctcggaa tgggcacccg agttcgccat ccacgacctg gattcacttg 480 atgegettge egeegeegae eeegeegaet tteeggaega categegeae tacetetgga 540 cctggaccga ccgctccgct ggctcgctgc gcgcccgcat gtttgccgcc aacttgggcg 600 tcaccgaaga cgaagcgacc ggtgccgcgg ccatccggat taccgattac ctcagccgtg 660 acctcaccat cacccaqqqc aaaqqatcqt tqatccacac cacctqqaqt cccqaqqqct 720 gggttcgggt agccggccga gttgtcagcg acggtgtggc acaactcgac tgacgtagag 780 ctcagcgctg ccgatgcaac acggcggcaa ggtgatcctg caggggttgc ccgaccqcqc 840 gcatctgcaa cgagtacgaa agctcgtcgc cgtcgatgcg gtaggaacgg tcaagggcgg 900 <210> 66

<211> 228

<212> PRT

<213> Mycobacterium tuberculosis

<400> 66

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Asn Phe Gly Asn Pro Leu Gly Val Ile Asn Ala Ser Lys Val Glu His 20 25 30

Arg Asp Arg Gln Gln Leu Ala Ala Gln Ser Gly Tyr Ser Glu Thr Ile 35 40 45

Phe Val Asp Leu Pro Ser Pro Gly Ser Thr Thr Ala His Ala Thr Ile 50 55 60

His Thr Pro Arg Thr Glu Ile Pro Phe Ala Gly His Pro Thr Val Gly 65 70 75 80

Ala Ser Trp Trp Leu Arg Glu Arg Gly Thr Pro Ile Asn Thr Leu Gln 85 90 95

Val Pro Ala Gly Ile Val Gln Val Ser Tyr His Gly Asp Leu Thr Ala 100 105 110

Ile Ser Ala Arg Ser Glu Trp Ala Pro Glu Phe Ala Ile His Asp Leu 115 120 125

Asp Ser Leu Asp Ala Leu Ala Ala Asp Pro Ala Asp Phe Pro Asp 130 135 140

Asp Ile Ala His Tyr Leu Trp Thr Trp Thr Asp Arg Ser Ala Gly Ser 145 150 155 160

Leu Arg Ala Arg Met Phe Ala Ala Asn Leu Gly Val Thr Glu Asp Glu
165 170 175

Ala Thr Gly Ala Ala Ile Arg Ile Thr Asp Tyr Leu Ser Arg Asp 180 185 190

Leu Thr Ile Thr Gln Gly Lys Gly Ser Leu Ile His Thr Thr Trp Ser 195 200 205

Pro Glu Gly Trp Val Arg Val Ala Gly Arg Val Val Ser Asp Gly Val 210 215 220

Ala Gln Leu Asp 225

<210> 67

<211> 500

<212> DNA

<213> Mycobacterium tuberculosis

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<210> 68

<211> 139

<212> PRT

<213> Mycobacterium tuberculosis

<400> 68

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Gly Ala Gly Ser Gly Pro Ala Met Gly Met Gly Gly Val Gly Gly Leu 20 25 30

Gly Gly Ala Gly Ser Gly Pro Ala Met Gly Met Gly Gly Val Gly Gly 35 40 45

Leu Asp Ala Ala Gly Ser Gly Glu Gly Gly Ser Pro Ala Ala Ile Gly 50 55 60

Ala Asp Thr Asn Arg Ser Asp Arg Ser Ser Asp Val Gly Gly Val 85 90 95

Trp Pro Leu Gly Phe Gly Arg Phe Ala Asp Ala Gly Ala Gly Gly Asn 100 105 110

Glu Ala Leu Gly Ser Lys Asn Gly Cys Ala Ala Ile Ser Ser Gly Ala 115 120 125

Ser Ile Pro Ser Cys Gly Arg Lys Ser Leu Ser 130 135

<210> 69

<211> 2050

<212> DNA

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accgagattt	cactcgactt	gctcgaccct	atctacaagc	gcaaggtcct	cgaattggcc	1980
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<210> 70

<211> 666

<212> PRT

<213> Mycobacterium tuberculosis

<400> 70

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Phe Pro Pro Ala Pro Ala Ser Ala Asn Leu Pro Lys Pro Asn Gly Gln
35 40 45

Thr Pro Pro Pro Thr Ser Asp Asp Leu Ser Glu Arg Phe Val Ser Ala 50 55 60

Pro Ile Ala Ala Gly Glu Pro Pro Ser Pro Glu Pro Ala Ala Ser Lys 85 90 95

Pro Pro Thr Pro Pro Met Pro Ile Ala Gly Pro Glu Pro Ala Pro Pro 100 105 110

Lys Pro Pro Thr Pro Pro Met Pro Ile Ala Gly Pro Glu Pro Ala Pro 115 120 125

Pro Lys Pro Pro Thr Pro Pro Met Pro Ile Ala Gly Pro Ala Pro Thr 130 135 140

Pro Thr Glu Ser Gln Leu Ala Pro Pro Arg Pro Pro Thr Pro Gln Thr 145 150 155 160

Pro Thr Gly Ala Pro Gln Gln Pro Glu Ser Pro Ala Pro His Val Pro 165 170 175

Ser	His	Gly	Pro 180	His	Gln	Pro	Arg	Arg 185	Thr	Ala	Pro	Ala	Pro 190	Pro	Trp
Ala	Lys	Met 195	Pro	Ile	Gly	Glu	Pro 200	Pro	Pro	Ala	Pro	Ser 205	Arg	Pro	Ser
Ala	Ser 210	Pro	Ala	Glu	Pro	Pro 215	Thr	Arg	Pro	Ala	Pro 220	Gln	His	Ser	Arg
Arg 225	Ala	Arg	Arg	Gly	His 230	Arg	Tyr	Arg	Thr	Asp 235	Thr	Glu	Arg	Asn	Val 240
Gly	Lys	Val	Ala	Thr 245	Gly	Pro	Ser	Ile	Gln 250	Ala	Arg	Leu	Arg	Ala 255	Glu
Glu	Ala	Ser	Gly 260	Ala	Gln	Leu	Ala	Pro 265	Gly	Thr	Glu	Pro	Ser 270	Pro	Ala
Pro	Leu	Gly 275	Gln	Pro	Arg	Ser	Tyr 280	Leu	Ala	Pro	Pro	Thr 285	Arg	Pro	Ala
Pro	Thr 290	Glu	Pro	Pro	Pro	Ser 295	Pro	Ser	Pro	Gln	Arg 300	Asn	Ser	Gly	Arg
Arg 305	Ala	Glu	Arg	Arg	Val 310	His	Pro	Asp	Leu	Ala 315	Ala	Gln	His	Ala	Ala 320
Ala	Gln	Pro	Asp	Ser 325	Ile	Thr	Ala	Ala	Thr 330	Thr	Gly	Gly	Arg	Arg 335	Arg
Lys	Arg	Ala	Ala 340	Pro	Asp	Leu	Asp	Ala 345	Thr	Gln	Lys	Ser	Leu 350	Arg	Pro
Ala	Ala	Lys 355	Gly	Pro	Lys	Val	Lys 360	Lys	Val	Lys	Pro	Gln 365	Lys	Pro	Lys
Ala	Thr 370	Lys	Pro	Pro	Lys	Val 375	Val	Ser	Gln	Arg	Gly 380	Trp	Arg	His	Trp
Val 385	His	Ala	Leu	Thr	Arg 390	Ile	Asn	Leu	Gly	Leu 395	Ser	Pro	Asp	Glu	Lys 400
Tyr	Glu	Leu	Asp	Leu 405	His	Ala	Arg	Val	Arg 410	Arg	Asn	Pro	Arg	Gly 415	Ser
Tyr	Gln	Ile	Ala 420	Val	Val	Gly	Leu	Lys 425	Gly	Gly	Ala	Gly	Lys 430	Thr	Thr
Leu	Thr	Ala 435	Ala	Leu	Gly	Ser	Thr 440	Leu	Ala	Gln	Val	Arg 445	Ala	Asp	Arg
Ile	Leu 450	Ala	Leu	Asp	Ala	Asp 455	Pro	Gly	Ala	Gly	Asn 460	Leu	Ala	Asp	Arg
Val 465	Gly	Arg	Gln	Ser	Gly 470	Ala	Thr	Ile	Ala	Asp 475	Val	Leu	Ala	Glu	Lys

Glu Leu Ser His Tyr Asn Asp Ile Arg Ala His Thr Ser Val Asn Ala 490 Val Asn Leu Glu Val Leu Pro Ala Pro Glu Tyr Ser Ser Ala Gln Arg Ala Leu Ser Asp Ala Asp Trp His Phe Ile Ala Asp Pro Ala Ser Arg 525 Phe Tyr Asn Leu Val Leu Ala Asp Cys Gly Ala Gly Phe Phe Asp Pro Leu Thr Arg Gly Val Leu Ser Thr Val Ser Gly Val Val Val Ala Ser Val Ser Ile Asp Gly Ala Gln Gln Ala Ser Val Ala Leu Asp Trp 565 Leu Arg Asn Asn Gly Tyr Gln Asp Leu Ala Ser Arg Ala Cys Val Val 585 Ile Asn His Ile Met Pro Gly Glu Pro Asn Val Ala Val Lys Asp Leu 600 Val Arg His Phe Glu Gln Gln Val Gln Pro Gly Arg Val Val Met 610 615 Pro Trp Asp Arg His Ile Ala Ala Gly Thr Glu Ile Ser Leu Asp Leu 630 Leu Asp Pro Ile Tyr Lys Arg Lys Val Leu Glu Leu Ala Ala Leu 645 Ser Asp Asp Phe Glu Arg Ala Gly Arg Arg 660 <210> 71 <211> 1890 <212> DNA Mycobacterium tuberculosis <213> <400> gcagcgatga ggaggagcgg cgccaacggc ccgcgccggc gacgatgcaa agcgcagcga 60 tgaggaggag cggcgcgcat gactgctgaa ccggaagtac ggacgctgcg cgaggttgtg 120 ctqqaccaqc tcqqcactqc tqaatcqcqt qcqtacaaqa tqtqqctqcc qccqttgacc 180 aatccggtcc cgctcaacga gctcatcgcc cgtgatcggc gacaacccct gcgatttgcc 240 ctqqqqatca tqqatqaacc qcqccqccat ctacaqqatq tqtqqqqcgt agacqtttcc 300 ggggccggcg gcaacatcgg tattgggggc gcacctcaaa ccgggaagtc gacgctactg 360 cagacgatgg tgatgtcggc cgccgccaca cactcaccgc gcaacgttca gttctattgc 420 480 ategacetag gtggcggcgg getgatetat etegaaaace tteeacacgt eggtggggta

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<210> 72

<211> 591

<212> PRT

<213> Mycobacterium tuberculosis

<400> 72

Met Thr Ala Glu Pro Glu Val Arg Thr Leu Arg Glu Val Val Leu Asp
1 5 10 15

Gln Leu Gly Thr Ala Glu Ser Arg Ala Tyr Lys Met Trp Leu Pro Pro 20 25 30

Leu Thr Asn Pro Val Pro Leu Asn Glu Leu Ile Ala Arg Asp Arg Arg 35 40 45

Gln Pro Leu Arg Phe Ala Leu Gly Ile Met Asp Glu Pro Arg Arg His 50 55 60

Leu Gln Asp Val Trp Gly Val Asp Val Ser Gly Ala Gly Gly Asn Ile 70 75 80

Gly Ile Gly Gly Ala Pro Gln Thr Gly Lys Ser Thr Leu Leu Gln Thr 85 90 95

Met Val Met Ser Ala Ala Ala Thr His Ser Pro Arg Asn Val Gln Phe 100 105 110

Tyr Cys Ile Asp Leu Gly Gly Gly Leu Ile Tyr Leu Glu Asn Leu 115 120 125

Pro His Val Gly Gly Val Ala Asn Arg Ser Glu Pro Asp Lys Val Asn 130 135 140

Arg Val Val Ala Glu Met Gln Ala Val Met Arg Gln Arg Glu Thr Thr
145 150 155 160

Phe Lys Glu His Arg Val Gly Ser Ile Gly Met Tyr Arg Gln Leu Arg 165 170 175

Asp Asp Pro Ser Gln Pro Val Ala Ser Asp Pro Tyr Gly Asp Val Phe
180 185 190

Leu Ile Ile Asp Gly Trp Pro Gly Phe Val Gly Glu Phe Pro Asp Leu 195 200 205

Glu Gly Gln Val Gln Asp Leu Ala Ala Gln Gly Leu Gly Phe Gly Val 210 215 220

His Val Ile Ile Ser Thr Pro Arg Trp Thr Glu Leu Lys Ser Arg Val 225 230 235 240

Arg Asp Tyr Leu Gly Thr Lys Ile Glu Phe Arg Leu Gly Asp Val Asn 245 250 255

Glu Thr Gln Ile Asp Arg Ile Thr Arg Glu Ile Pro Ala Asn Arg Pro
260 265 270

Gly Arg Ala Val Ser Met Glu Lys His His Leu Met Ile Gly Val Pro 275 280 285

Arg Phe Asp Gly Val His Ser Ala Asp Asn Leu Val Glu Ala Ile Thr 290 295 300

Ala Gly Val Thr Gln Ile Ala Ser Gln His Thr Glu Gln Ala Pro Pro 305 310 315 Val Arg Val Leu Pro Glu Arg Ile His Leu His Glu Leu Asp Pro Asn 330 Pro Pro Gly Pro Glu Ser Asp Tyr Arg Thr Arg Trp Glu Ile Pro Ile Gly Leu Arg Glu Thr Asp Leu Thr Pro Ala His Cys His Met His Thr Asn Pro His Leu Leu Ile Phe Gly Ala Ala Lys Ser Gly Lys Thr Thr Ile Ala His Ala Ile Ala Arg Ala Ile Cys Ala Arg Asn Ser Pro Gln 390 Gln Val Arg Phe Met Leu Ala Asp Tyr Arg Ser Gly Leu Leu Asp Ala 410 Val Pro Asp Thr His Leu Leu Gly Ala Gly Ala Ile Asn Arg Asn Ser Ala Ser Leu Asp Glu Ala Ala Gln Ala Leu Ala Val Asn Leu Lys Lys 440 435 445 Arg Leu Pro Pro Thr Asp Leu Thr Thr Ala Gln Leu Arg Ser Arg Ser 455 Trp Trp Ser Gly Phe Asp Val Val Leu Leu Val Asp Asp Trp His Met 465 470 480 Ile Val Gly Ala Ala Gly Gly Met Pro Pro Met Ala Pro Leu Ala Pro 490 Leu Leu Pro Ala Ala Ala Asp Ile Gly Leu His Ile Ile Val Thr Cys 505 Gln Met Ser Gln Ala Tyr Lys Ala Thr Met Asp Lys Phe Val Gly Ala 515 520 Ala Phe Gly Ser Gly Ala Pro Thr Met Phe Leu Ser Gly Glu Lys Gln 535 Glu Phe Pro Ser Ser Glu Phe Lys Val Lys Arg Arg Pro Pro Gly Gln 545 555 550 560 Ala Phe Leu Val Ser Pro Asp Gly Lys Glu Val Ile Gln Ala Pro Tyr 565 570 Ile Glu Pro Pro Glu Glu Val Phe Ala Ala Pro Pro Ser Ala Gly 585

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acatgatccg atcgctgccg acattggcac gcaagtgagc gacaacgctc tgcacggcgt
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Phe Ala	
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gctgggttgc cgccgccggt gccgcccacc tggctgaaca acgacgtcac ctgctgcagc	240
ggctgggtca gctgctgcat cgggccgctc atctcaccca gttggccgag ggtctgggta	300
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Ala Gly Leu Pro Pro Pro Val Pro Pro Thr Trp Leu Asn Asn Asp Val

50 55 60

Thr Cys Cys Ser Gly Trp Val Ser Cys Cys Ile Gly Pro Leu Ile Ser 65 70 75 80

Pro Ser Trp Pro Arg Val Trp Val Ala Ala Gly Gly Asn Trp Pro Thr
85 90 95

Gly Val Glu Leu Pro Gly Glu Gly Ile Pro Lys Ile Gly Phe Val Val 100 105 110

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Asp Lys Ala Let	ı Ala Ala Ala T 70	Thr Pro Met Val 75	Val Trp Leu Gln Thr 80	
Ala Ser Thr Gli	n Ala Lys Thr A 85	Arg Ala Met Gln 90	Ala Thr Ala Gln Ala 95	
Ala Ala Tyr Thi		Ala Thr Thr Pro 105	Ser Leu Pro Glu Ile 110	
Ala Ala Asn His 115		Ala Val Leu Thr 120	Ala Thr Asn Phe Phe 125	
Gly Ile Asn Thi	Tile Pro Ile A	Ala Leu Thr Glu	Met Asp Tyr Phe Ile 140	
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Thr Ala Val Ası	n Thr Leu Phe G 165	Glu Lys Leu Glu 170	Pro Met Ala Ser Ile 175	
Leu Asp Pro Gly		Ser Thr Thr Asn 185	Pro Ile Phe Gly Met 190	
Pro Ser Pro Gly		Pro Val Gly Gln 200	Leu Pro Pro Ala Ala 205	
Thr Gln Thr Let	ı Gly Gln Leu G 215	Gly Glu Met Ser	Gly Pro Met Gln Gln 220	

Leu Thr Gln Pro Leu Gln Gln Val Thr Ser Leu Phe Ser Gln Val Gly

235 240 225 230 Gly Thr Gly Gly Gly Asn Pro Ala Asp Glu Glu Ala Ala Gln Met Gly 245 250 Leu Leu Gly Thr Ser Pro Leu Ser Asn His Pro Leu Ala Gly Gly Ser 265 Gly Pro Ser Ala Gly Ala Gly Leu Leu Arg Ala Glu Ser Leu Pro Gly 280 Ala Gly Gly Ser Leu Thr Arg Thr Pro Leu Met Ser Gln Leu Ile Glu 295 Lys Pro Val Ala Pro Ser Val Met Pro Ala Ala Ala Gly Ser Ser 310 315 Ala Thr Gly Gly Ala Ala Pro Val Gly Ala Gly Ala Met Gly Gln Gly 325 Ala Gln Ser Gly Gly Ser Thr Arg Pro Gly Leu Val Ala Pro Ala Pro Leu Ala Gln Glu Arg Glu Glu Asp Asp Glu Asp Asp Trp Asp Glu Glu Asp Asp Trp 370 <210> 93 <211> 1000 <212> DNA <213> Mycobacterium tuberculosis gacgcgacac agaaatcctt aaggccggcg gccaaggggc cgaaggtgaa gaaggtgaag 60 ccccagaaac cgaaggccac gaagccgccc aaagtggtgt cgcagcgcgg ctggcgacat 120 tgggtgcatg cgttgacgcg aatcaacctg ggcctgtcac ccgacgagaa gtacgagctg 180 gacctgcacg ctcgagtccg ccgcaatccc cgcgggtcgt atcagatcgc cgtcgtcggt 240 300 ctcaaaggtg gggctggcaa aaccacgctg acagcagcgt tggggtcgac gttggctcag gtgcgggccg accggatcct ggctctagac gcggatccag gcgccggaaa cctcgccgat 360 cqqqtaqqqc gacaatcqqq cqcqaccatc gctgatqtqc ttgcaqaaaa agaqctqtcq 420 cactacaacg acatecgege acacactage gteaatgegg teaatetgga agtgetgeeg 480 gcaccggaat acagetegge gcagegegeg etcagegaeg ecgaetggea tttcategee 540 gatcctgcgt cgaggtttta caacctcgtc ttggctgatt gtggggccgg cttcttcgac 600

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660

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Gly Leu Lys Gly Gly Ala Gly Lys Thr Thr Leu Thr Ala Ala Leu Gly 65 70 75 80

Ser Thr Leu Ala Gln Val Arg Ala Asp Arg Ile Leu Ala Leu Asp Ala 85 90 95

Asp Pro Gly Ala Gly Asn Leu Ala Asp Arg Val Gly Arg Gln Ser Gly
100 105 110

Ala Thr Ile Ala Asp Val Leu Ala Glu Lys Glu Leu Ser His Tyr Asn 115 120 125

Asp Ile Arg Ala His Thr Ser Val Asn Ala Val Asn Leu Glu Val Leu 130 135 140

Pro Ala Pro Glu Tyr Ser Ser Ala Gln Arg Ala Leu Ser Asp Ala Asp 145 150 155 160

Trp His Phe Ile Ala Asp Pro Ala Ser Arg Phe Tyr Asn Leu Val Leu 165 170 175

Ala Asp Cys Gly Ala Gly Phe Phe Asp Pro Leu Thr Arg Gly Val Leu 180 185 190

Ser Thr Val Ser Gly Val Val Val Ala Ser Val Ser Ile Asp Gly

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Val Ser
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<213> Mycobacterium tuberculosis

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010	124	
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35 3		
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gaagagtgtc	tcggccagcg	gcgatacctt	gggtgccgtc	atcagcgacc	tggaggccaa	180
ctattcgggc	atttccgagc	gcctgatgga	cccgtcttcc	ccaggtaagt	tgcaccgctt	240
cgtgaacatc	tacgtcaacg	acgaggacgt	gcggttctcc	ggcggcttgg	ccaccgcgat	300
cgctgacggt	gactcggtca	ccatcctccc	cgccgtggcc	ggtgggtgag	cggagcacat	360
gacacgatac	gactcgctgt	tgcaggcctt	gggcaacacg	ccgctggttg	gcctgcagcg	420
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<211> 93

<212> PRT

<213> Mycobacterium tuberculosis

<400> 141

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Gly Gln Lys Ser Val Ser Ala Ser Gly Asp Thr Leu Gly Ala Val Ile 20 25 30

Ser Asp Leu Glu Ala Asn Tyr Ser Gly Ile Ser Glu Arg Leu Met Asp 35 40 45

Pro Ser Ser Pro Gly Lys Leu His Arg Phe Val Asn Ile Tyr Val Asn 50 55 60

Asp Glu Asp Val Arg Phe Ser Gly Gly Leu Ala Thr Ala Ile Ala Asp 65 70 75 80

Gly Asp Ser Val Thr Ile Leu Pro Ala Val Ala Gly Gly 85 90

<210> 142

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<212> DNA

<213> Mycobacterium tuberculosis '

<400> 142

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atgcgggaca	tggcgggccg	ttttgaggtg	cacgcccaga	cggtggagga	cgaggctcgc	180
cggatgtggg	cgtccgcgca	aaacatctcg	ggcgcgggct	ggagtggcat	ggccgaggcg	240
acctcgctag	acaccatggc	ccagatgaat	caggcgtttc	gcaacatcgt	gaacatgctg	300
cacggggtgc	gtgacgggct	ggttcgcgac	gccaacaact	acgagcagca	agagcaggcc	360
tcccagcaga	tcctcagcag	ctaacgtcag	ccgctgcagc	acaatacttt	tacaagcgaa	420
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<210> 143

<211> 98

<212> PRT

<213> Mycobacterium tuberculosis

<400> 143

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20 25 30

Met Trp Ala Ser Ala Gln Asn Ile Ser Gly Ala Gly Trp Ser Gly Met 35 40 45

Ala Glu Ala Thr Ser Leu Asp Thr Met Ala Gln Met Asn Gln Ala Phe 50 55 60

Arg Asn Ile Val Asn Met Leu His Gly Val Arg Asp Gly Leu Val Arg 65 70 75 80

Asp Ala Asn Asn Tyr Glu Gln Glu Gln Ala Ser Gln Gln Ile Leu 85 90 95

Ser Ser

<210> 144

<211> 940

<212> DNA

<213> Mycobacterium tuberculosis

<400> 144

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gaactgggte gacetteaga ceaeeggtea gteegeegee aaaaagttet acacategtt 180
gtteggetgg ggttacgaeg acaaeeeggt eeeeggagge ggtggggtet atteeatgge 240

cacqctqaac qqcqaaqccq tqqccqccat cqcaccqatq cccccqqqtq caccqgaggg 300 qatqccqccq atctqqaaca cctatatcgc ggtggacgac gtcgatgcgg tggtggacaa 360 qqtqqtgccc gggggcgggc aggtgatgat gccggccttc gacatcggcg atgccggccg 420 qatqtcqttc atcaccqatc cgaccggcgc tgccgtgggc ctatggcagg ccaatcggca 480 categgageg aegttggtea aegagaeggg caegeteate tggaaegaae tgeteaegga 540 caaqccggat ttggcgctag cgttctacga ggctgtggtt ggcctcaccc actcgagcat 600 ggagataget gegggeeaga actategggt geteaaggee ggegaegegg aagteggegg 660 ctqtatqqaa ccqccqatqc ccqqcqtgcc gaatcattgg cacgtctact ttgcggtgga 720 tqacqccqac qccacqqcqq ccaaaqccqc cgcagcgggc ggccaggtca ttgcggaacc 780 ggctgacatt ccgtcggtgg gccggttcgc cgtgttgtcc gatccgcagg gcgcgatctt 840 caqtqtqttq aaqcccgcac cgcagcaata gggagcatcc cgggcaggcc cgccggccgg 900 940 cagattegga gaatgetaga agetgeegee ggegeegeeg

<210> 145

<211> 261

<212> PRT

<213> Mycobacterium tuberculosis

<400> 145

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Leu Gln Thr Thr Asp Gln Ser Ala Ala Lys Lys Phe Tyr Thr Ser Leu 20 25 30

Phe Gly Trp Gly Tyr Asp Asp Asn Pro Val Pro Gly Gly Gly Val 35 40 45

Tyr Ser Met Ala Thr Leu Asn Gly Glu Ala Val Ala Ala Ile Ala Pro 50 55 60

Met Pro Pro Gly Ala Pro Glu Gly Met Pro Pro Ile Trp Asn Thr Tyr 65 70 75 80

Ile Ala Val Asp Asp Val Asp Ala Val Val Asp Lys Val Val Pro Gly
85 90 95

Gly Gly Gln Val Met Met Pro Ala Phe Asp Ile Gly Asp Ala Gly Arg
100 105 110

Met Ser Phe Ile Thr Asp Pro Thr Gly Ala Ala Val Gly Leu Trp Gln
115 120 125

Ala	Asn 130	Arg	His	Ile	Gly	Ala 135	Thr	Leu	Val	Asn	Glu 140	Thr	Gly	Thr	Leu	
Ile 145	Trp	Asn	Glu	Leu	Leu 150	Thr	Asp	Lys	Pro	Asp 155	Leu	Ala	Leu	Ala	Phe 160	
Tyr	Glu	Ala	Val	Val 165	Gly	Leu	Thr	His	Ser 170	Ser	Met	Glu	Ile	Ala 175	Ala	
Gly	Gln	Asn	Tyr 180	Arg	Val	Leu	Lys	Ala 185	Gly	Asp	Ala	Glu	Val 190	Gly	Gly	
Cys	Met	Glu 195	Pro	Pro	Met	Pro	Gly 200	Val	Pro	Asn	His	Trp 205	His	Val	Tyr	
Phe	Ala 210	Val	Asp	Asp	Ala	Asp 215	Ala	Thr	Ala	Ala	Lys 220	Ala	Ala	Ala	Ala	
Gly 225	Gly	Gln	Val	Ile	Ala 230	Glu	Pro	Ala	Asp	Ile 235	Pro	Ser	Val	Gly	Arg 240	
Phe	Ala	Val	Leu	Ser 245	Asp	Pro	Gln	Gly	Ala 250	Ile	Phe	Ser	Val	Leu 255	Lys	
Pro	Ala	Pro	Gln 260	Gln												
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ggai	gtat	tt g	gtcca	acta	ca c	ggaga	atcca	a ggg	gaac	gggc	ttc	cgca	ccc 1	ttgaa	agaaaa	180
cca	gaagg	gtc g	gagti	cga	ga to	egge	caca	g cc	ctaaç	gggc	ccc	cagg	cca o	ccgga	agtccg	240
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Phe	Ile	Ala	Pro	Glu	Asp	Gly	Ser	Ala	Asp	Val	Phe	Val	His	Tyr	Thr	

Glu Ile Gln Gly Thr Gly Phe Arg Thr Leu Glu Glu Asn Gln Lys Val 35 40 45

Glu Phe Glu Ile Gly His Ser Pro Lys Gly Pro Gln Ala Thr Gly Val 50 55 60

Arg Ser Leu 65

<210> 148

<211> 540

<212> DNA

<213> Mycobacterium tuberculosis

<400> 148

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<210> 149

<211> 129

<212> PRT

<213> Mycobacterium tuberculosis

<400> 149

Met Asn Leu Arg Arg His Gln Thr Leu Thr Leu Arg Leu Leu Ala Ala 1 5 10 15

Ser Ala Gly Ile Leu Ser Ala Ala Ala Phe Ala Ala Pro Ala Gln Ala 20 25 30

Asn Pro Val Asp Asp Ala Phe Ile Ala Ala Leu Asn Asn Ala Gly Val 35 40 45

Asn Tyr Gly Asp Pro Val Asp Ala Lys Ala Leu Gly Gln Ser Val Cys
50 55 60

Pro Ile Leu Ala Glu Pro Gly Gly Ser Phe Asn Thr Ala Val Ala Ser 65 70 75 80

Val Val Ala Arg Ala Gln Gly Met Ser Gln Asp Met Ala Gln Thr Phe

85 90 95

Thr Ser Ile Ala Ile Ser Met Tyr Cys Pro Ser Val Met Ala Asp Val 100 105 110

Ala Ser Gly Asn Leu Pro Ala Leu Pro Asp Met Pro Gly Leu Pro Gly 115 120 125

Ser

	$\overline{}$	-	^		-	15	^
<	Z.	- 1	u	' >		רו	u

<211> 400

<212> DNA

<213> Mycobacterium tuberculosis

<400> 150

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<211> 110

<212> PRT

<213> Mycobacterium tuberculosis

<400> 151

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Met Ser Leu Thr Val Gly Ala Gly Val Ala Ser Ala Asp Pro Val Asp 20 25 30

Ala Val Ile Asn Thr Thr Cys Asn Tyr Gly Gln Val Val Ala Ala Leu 35 40 45

Asn Ala Thr Asp Pro Gly Ala Ala Ala Gln Phe Asn Ala Ser Pro Val 50 55 60

Ala Gln Ser Tyr Leu Arg Asn Phe Leu Ala Ala Pro Pro Pro Gln Arg
65 70 75 80

Ala Ala Met Ala Ala Gln Leu Gln Ala Val Pro Gly Ala Ala Gln Tyr
85 90 95

Ile Gly Leu Val Glu Ser Val Ala Gly Ser Cys Asn Asn Tyr 100 105 110

<210> 152

<211> 990

<212> DNA

<213> Mycobacterium tuberculosis

<400> 152

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<210> 153

<211> 266

<212> PRT

<213> Mycobacterium tuberculosis

<400> 153

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Glu Arg Lys Leu Thr Asp Gly Asp Phe Thr Leu Asp Arg Glu Ala Ala 20 25 30

Asp	Ala	Val	Leu	Asp	Glu	Ile	Asn	Glu	Arg	Ala	Val	Glu	Glu	Ala	Leu
		35					40					45			

Gln Ile Arg Glu Lys Glu Ala Ala Asp Gly Ile Glu Gly Ser Val Thr 50 55 60

Val Leu Thr Ala Gly Pro Glu Arg Ala Thr Glu Ala Ile Arg Lys Ala 65 70 75 80

Leu Ser Met Gly Ala Asp Lys Ala Val His Leu Lys Asp Asp Gly Met 85 90 95

His Gly Ser Asp Val Ile Gln Thr Gly Trp Ala Leu Ala Arg Ala Leu 100 105 110

Gly Thr Ile Glu Gly Thr Glu Leu Val Ile Ala Gly Asn Glu Ser Thr 115 120 125

Asp Gly Val Gly Gly Ala Val Pro Ala Ile Ile Ala Glu Tyr Leu Gly 130 135 140

Leu Pro Gln Leu Thr His Leu Arg Lys Val Ser Ile Glu Gly Gly Lys
145 150 155 160

Ile Thr Gly Glu Arg Glu Thr Asp Glu Gly Val Phe Thr Leu Glu Ala 165 170 175

Thr Leu Pro Ala Val Ile Ser Val Asn Glu Lys Ile Asn Glu Pro Arg 180 185 190

Phe Pro Ser Phe Lys Gly Ile Met Ala Ala Lys Lys Glu Val Thr

Val Leu Thr Leu Ala Glu Ile Gly Val Glu Ser Asp Glu Val Gly Leu 210 215 220

Ala Asn Ala Gly Ser Thr Val Leu Ala Ser Thr Pro Lys Pro Ala Lys 225 230 235 240

Thr Ala Gly Glu Lys Val Thr Asp Glu Gly Glu Gly Gly Asn Gln Ile 245 250 255

Val Gln Tyr Leu Val Ala Gln Lys Ile Ile 260 265

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<212> DNA

<213> Mycobacterium tuberculosis

<400> 154

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<210> 155

<211> 35

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	ggge councingery regengings	
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55-		
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-2137	11 COMOCOLIUM CUMOLOULOULO	
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tctccc	gggg gtaactcaga gcgagcggac	30
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Pro Gln Gly Thr Val Lys Trp Phe Asn Ala Glu Lys Gly Phe Gly
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               5
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Thr Arg Phe Met Thr Asp Pro His Ala Met Arg Asp Met Ala Gly
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Asn Thr Arg Leu Trp Val Tyr Cys Gly Asn Gly Thr Pro Asn Glu Leu

Trp Glu Arg Asn Asp Pro Thr Gln Gln Ile Pro Lys Leu Val Ala Asn

215

225	230	235	240

Gly Gly Ala Asn Ile Pro Ala Glu Phe Leu Glu Asn Phe Val Arg Ser 245 250 255

Ser Asn Leu Lys Phe Gln Asp Ala Tyr Asn Ala Ala Gly Gly His Asn 260 265 270

Ala Val Phe Asn Phe Pro Pro Asn Gly Thr His Ser Trp Glu Tyr Trp 275 280 285

Gly Ala Gln Leu Asn Ala Met Lys Gly Asp Leu Gln Ser Ser Leu Gly 290 295 300

Ala Gly Lys Leu Ala Met Thr Glu Gln Gln Trp Asn Phe Ala Gly Ile 305 310 315 320

Glu Ala Ala Ala Ser Ala Ile Gln Gly Asn Val Thr Ser Ile His Ser 325 330 335

Leu Leu Asp Glu Gly Lys Gln Ser Leu Thr Lys Leu Ala Ala Ala Trp
340 345 350

Gly Gly Ser Gly Ser Glu Ala Tyr Gln Gly Val Gln Gln Lys Trp Asp 355 360 365

Ala Thr Ala Thr Glu Leu Asn Asn Ala Leu Gln Asn Leu Ala Arg Thr 370 375 380

Ile Ser Glu Ala Gly Gln Ala Met Ala Ser Thr Glu Gly Asn Val Thr 385 390 395 400

Gly Met Phe Ala

<210> 173

<211> 403

<212> PRT

<213> Mycobacterium tuberculosis

<400> 173

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Glu Ala Ala Ser Ala Ile Gln Gly Asn Val Thr Ser Ile His Ser 35 40 45

Leu Leu Asp Glu Gly Lys Gln Ser Leu Thr Lys Leu Ala Ala Ala Trp 50 55 60

Gly Gly Ser Gly Ser Glu Ala Tyr Gln Gly Val Gln Gln Lys Trp Asp
65 70 75 80

Ala	Thr	Ala	Thr	Glu 85	Leu	Asn	Asn	Ala	Leu 90	Gln	Asn	Leu	Ala	Arg 95	Thr
Ile	Ser	Glu	Ala 100	Gly	Gln	Ala	Met	Ala 105	Ser	Thr	Glu	Gly	Asn 110	Val	Thr
Gly	Met	Phe 115	Ala	Lys	Leu	Phe	Ser 120	Arg	Pro	Gly	Leu	Pro 125	Val	Glu	Tyr
Leu	Gln 130	Val	Pro	Ser	Pro	Ser 135	Met	Gly	Arg	Asp	Ile 140	Lys	Val	Gln	Phe
Gln 145	Ser	Gly	Gly	Asn	Asn 150	Ser	Pro	Ala	Val	Tyr 155	Leu	Leu	Asp	Gly	Leu 160
Arg	Ala	Gln	Asp	Asp 165	Tyr	Asn	Gly	Trp	Asp 170	Ile	Asn	Thr	Pro	Ala 175	Phe
Glu	Trp	Tyr	Tyr 180	Gln	Ser	Gly	Leu	Ser 185	Ile	Val	Met	Pro	Val 190	Gly	Gly
Gln	Ser	Ser 195	Phe	Tyr	Ser	Asp	Trp 200	Tyr	Ser	Pro	Ala	Cys 205	Gly	Lys	Ala
Gly	Cys 210	Gln	Thr	Tyr	Lys	Trp 215	Glu	Thr	Phe	Leu	Thr 220	Ser	Glu	Leu	Pro
Gln 225	Trp	Leu	Ser	Ala	Asn 230	Arg	Ala	Val	Lys	Pro 235	Thr	Gly	Ser	Ala	Ala 240
Ile	Gly	Leu	Ser	Met 245	Ala	Gly	Ser	Ser	Ala 250	Met	Ile	Leu	Ala	Ala 255	Tyr
His	Pro	Gln	Gln 260	Phe	Ile	Tyr	Ala	Gly 265	Ser	Leu	Ser	Ala	Leu 270	Leu	Asp
Pro	Ser	Gln 275	Gly	Met	Gly	Pro	Ser 280	Leu	Ile	Gly	Leu	Ala 285	Met	Gly	Asp
Ala	Gly 290	Gly	Tyr	Lys	Ala	Ala 295	Asp	Met	Trp	Gly	Pro 300	Ser	Ser	Asp	Pro
Ala 305	Trp	Glu	Arg	Asn	Asp 310	Pro	Thr	Gln	Gln	Ile 315	Pro	Lys	Leu	Val	Ala 320
Asn	Asn	Thr	Arg	Leu 325	Trp	Val	Tyr	Cys	Gly 330	Asn	Gly	Thr	Pro	Asn 335	Glu
Leu	Gly	Gly	Ala 340	Asn	Ile	Pro	Ala	Glu 345	Phe	Leu	Glu	Asn	Phe 350	Val	Arg
Ser	Ser	Asn 355	Leu	Lys	Phe	Gln	Asp 360	Ala	Tyr	Asn	Ala	Ala 365	Gly	Gly	His
Asn	Ala	Val	Phe	Asn	Phe	Pro	Pro	Asn	Gly	Thr	His	Ser	Trp	Glu	Tyr

Trp Gly Ala Gln Leu Asn Ala Met Lys Gly Asp Leu Gln Ser Ser L 385 390 395 4	eu 100
Gly Ala Gly	
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agtgcttggc agggtgatac cgggatcacg tatcagggct ggcagaccca gtggaa	accag
gccctagagg atctggtgcg ggcctatcag tcgatgtctg gcacccatga gtccaa	icacc :
atggcgatgt tggctcgaga tggggccgaa gccgccaagt ggggcggcta g	:
<pre><400> 175 Met Ser Gln Ile Met Tyr Asn Tyr Pro Ala Met Met Ala His Ala G 1</pre>	Bly
	Sly
Asp Met Ala Gly Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Asp I 20 25 30	Ile
Ala Ser Glu Gln Ala Val Leu Ser Ser Ala Trp Gln Gly Asp Thr G 35 40 45	Bly
Ile Thr Tyr Gln Gly Trp Gln Thr Gln Trp Asn Gln Ala Leu Glu A 50 55 60	Asp
Leu Val Arg Ala Tyr Gln Ser Met Ser Gly Thr His Glu Ser Asn T	Thr 30
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gccatggagg ctctcgcgcg ggcctaccgt cggtgccggc gagcactacg ccagatcggg	240
gtgctggaaa ggccggtagg cgattcgtca gactgcggaa cgattagggt ggggtcgttc	300
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Ala Ser Glu Arg Thr Ala Pro Ser Arg Ala Cys Gln Gly Asp Leu Gly 35 40 45	
Met Ser His Gln Asp Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Ala 50 55 60	
Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu Arg Gln Ile Gly 70 75 80	
Val Leu Glu Arg Pro Val Gly Asp Ser Ser Asp Cys Gly Thr Ile Arg 85 90 95	
Val Gly Ser Phe Arg Gly Arg Trp Leu Asp Pro Arg His Ala Gly Pro 100 105 110	
Ala Thr Ala Ala Asp Ala Gly Asp 115 120	
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tegggegegg getggagtgg catggeegag gegaeetege tagacaceat gaeecagatg	180
aatcaggegt ttegeaacat egtgaacatg etgeaegggg tgegtgaegg getggttege	240

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<400>	179															
Met Ala	a Ser	Arg	Phe 5	Met	Thr	Asp	Pro	His 10	Ala	Met	Arg	Asp	Met 15	Ala		
Gly Arg	g Phe	Glu 20	Val	His	Ala	Gln	Thr 25	Val	Glu	Asp	Glu	Ala 30	Arg	Arg		
Met Tr	Ala 35	Ser	Ala	Gln	Asn	Ile 40	Ser	Gly	Ala	Gly	Trp 45	Ser	Gly	Met		
Ala Glu 50	ı Ala	Thr	Ser	Leu	Asp 55	Thr	Met	Thr	Gln	Met 60	Asn	Gln	Ala	Phe		
Arg Ası 65	ı Ile	Val	Asn	Met 70	Leu	His	Gly	Val	Arg 75	Asp	Gly	Leu	Val	Arg 80		
Asp Ala	a Asn	Asn	Tyr 85	Glu	Gln	Gln	Glu	Gln 90	Ala	Ser	Gln	Gln	Ile 95	Leu		
Ser Sei	2															
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tccggt															180	ı
aatcag															240)
gacgcca															297	
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Met Ala Ser Arg Phe Met Thr Asp Pro His Ala Met Arg Asp Met Ala

10

15

Gly	Arg	Phe	Glu 20	Val	His	Ala	Gln	Thr 25	Val	Glu	Asp	Glu	Ala 30	Arg	Arg	
Met	Trp	Ala 35	Ser	Ala	Gln	Asn	Ile 40	Ser	Gly	Ala	Gly	Trp 45	Ser	Gly	Met	
Ala	Glu 50	Ala	Thr	Ser	Leu	Asp 55	Thr	Met	Ala	Gln	Met 60	Asn	Gln	Ala	Phe	
Arg 65	Asn	Ile	Val	Asn	Met 70	Leu	His	Gly	Val	Arg 75	Asp	Gly	Leu	Val	Arg 80	
Asp	Ala	Asn	Asn	Tyr 85	Glu	Gln	Gln	Glu	Gln 90	Ala	Ser	Gln	Gln	Ile 95	Leu	
Ser	Ser															
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<400		182					.~~.		*	.~~~	2021	- ~~~		aaet	-++	60
	-		-				_								ttgag	
gtgo	cacgo	CCC a	agaco	gtgg	ga go	gacga	aggct	cg(ccgga	atgt	ggg	cgtco	ege g	gcaaa	aacatt	120
tcc	gtg	egg g	gctgo	gagto	g ca	atggo	ccgaç	g gcg	gacct	cgc	taga	acaco	cat g	gacct	agatg	180
aato	aggo	egt t	tcg	caaca	at co	gtgaa	acato	gct	gcaco	9999	tgcg	gtgad	egg g	gctgg	gttcgc	240
gac	gccaa	aca a	actac	gaad	a go	caaga	agcag	g gco	ctcc	cagc	agat	ccts	gag d	cagct	ag	297
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Met	Trp	Ala 35	Ser	Ala	Gln	Asn	Ile 40	Ser	Gly	Ala	Gly	Trp 45	Ser	Gly	Met	
Ala	Glu 50	Ala	Thr	Ser	Leu	Asp 55	Thr	Met	Thr	Gln	Met 60	Asn	Gln	Ala	Phe	
Arg 65	Asn	Ile	Val	Asn	Met 70	Leu	His	Gly	Val	Arg 75	Asp	Gly	Leu	Val	Arg 80	

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Ser Ser	
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teeggegegg getggagtgg catggeegag gegacetege tagacaceat gacecagatg 1	80
aatcaggcgt ttcgcaacat cgtgaacatg ctgcacgggg tgcgtgacgg gctggttcgc 2	40
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Ala Glu Ala Thr Ser Leu Asp Thr Met Thr Gln Met Asn Gln Ala Phe 50 55 60	
Arg Asn Ile Val Asn Met Leu His Gly Val Arg Asp Gly Leu Val Arg 65 70 75 80	
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atcgaa	ecct gctgaccgag aggacttgtg atg tcg caa atc atg tac aac tac Met Ser Gln Ile Met Tyr Asn Tyr 1 5	114
	g atg ttg ggt cac gcc ggg gat atg gcc gga tat gcc ggc acg a Met Leu Gly His Ala Gly Asp Met Ala Gly Tyr Ala Gly Thr 15 20	162
	g agc ttg ggt gcc gag atc gcc gtg gag cag gcc gcg ttg cag n Ser Leu Gly Ala Glu Ile Ala Val Glu Gln Ala Ala Leu Gln 30 35 40	210
	g tgg cag ggc gat acc ggg atc acg tat cag gcg tgg cag gca a Trp Gln Gly Asp Thr Gly Ile Thr Tyr Gln Ala Trp Gln Ala 45 50 55	258
	g aac cag gcc atg gaa gat ttg gtg cgg gcc tat cat gcg atg o Asn Gln Ala Met Glu Asp Leu Val Arg Ala Tyr His Ala Met 60 65 70	306
	e acc cat gaa gcc aac acc atg gcg atg atg gcc cgc gac acc r Thr His Glu Ala Asn Thr Met Ala Met Met Ala Arg Asp Thr 75 80 85	354
_	a gcc gcc aaa tgg ggc ggc tag 1 Ala Ala Lys Trp Gly Gly 95	381
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Met Ser Gln Ile Met Tyr Asn Tyr Pro Ala Met Leu Gly His Ala Gly Asp Met Ala Gly Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Glu Ile 25 Ala Val Glu Gln Ala Ala Leu Gln Ser Ala Trp Gln Gly Asp Thr Gly 40 Ile Thr Tyr Gln Ala Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Asp 55 Leu Val Arq Ala Tyr His Ala Met Ser Ser Thr His Glu Ala Asn Thr 70 65 Met Ala Met Met Ala Arg Asp Thr Ala Glu Ala Ala Lys Trp Gly Gly <210> 196 <211> 363 <212> DNA <213> Mycobacterium tuberculosis <220> <221> CDS <222> (1)..(360) <400> 196 gtg teg cag agt atg tac age tac eeg geg atg aeg gee aat gte gga 48 Val Ser Gln Ser Met Tyr Ser Tyr Pro Ala Met Thr Ala Asn Val Gly gac atg gcc ggt tat acg ggc acg acg cag agc ttg ggg gcc gat atc 96 Asp Met Ala Gly Tyr Thr Gly Thr Thr Gln Ser Leu Gly Ala Asp Ile gcc agt gag cgc acc gcg ccg tcg cgt gct tgc caa ggt gat ctc ggg 144 Ala Ser Glu Arg Thr Ala Pro Ser Arg Ala Cys Gln Gly Asp Leu Gly atq aqt cat caq qac tqq caq qcc cag tgg aat cag gcc atg gag gct 192 Met Ser His Gln Asp Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Ala 50 55 ctc gcg cgg gcc tac cgt cgg tgc cgg cga gca cta cgc cag atc ggg 240 Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu Arg Gln Ile Gly 70 75 80 65 gtg ctg gaa agg ccg gta ggc gat tcg tca gac tgc gga acg att agg 288

Val Leu Glu Arg Pro Val Gly Asp Ser Ser Asp Cys Gly Thr Ile Arg

336

363

gtg ggg tcg ttc cgg ggt cgg tgg ctg gac ccg cgc cat gcg ggt cca Val Gly Ser Phe Arg Gly Arg Trp Leu Asp Pro Arg His Ala Gly Pro 100 105 110

gcc acg gcc gcc ggc gga gac taa Ala Thr Ala Ala Asp Ala Gly Asp 115 120

<210> 197
<211> 120
<212> PRT

<213> Mycobacterium tuberculosis

<400> 197

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1 5 10 15

Asp Met Ala Gly Tyr Thr Gly Thr Thr Gln Ser Leu Gly Ala Asp Ile 20 25 30

Ala Ser Glu Arg Thr Ala Pro Ser Arg Ala Cys Gln Gly Asp Leu Gly 35 40 45

Met Ser His Gln Asp Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Ala 50 55 60

Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu Arg Gln Ile Gly 65 70 75 80

Val Leu Glu Arg Pro Val Gly Asp Ser Ser Asp Cys Gly Thr Ile Arg 85 90 95

Val Gly Ser Phe Arg Gly Arg Trp Leu Asp Pro Arg His Ala Gly Pro 100 105 110

Ala Thr Ala Ala Asp Ala Gly Asp 115 120

<210> 198

<211> 291

<212> DNA

<213> Mycobacterium tuberculosis

<220>

<221> CDS

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Leu Val Arg Ala Tyr Gln Ser Met Ser Gly Thr His Glu Ser Asn Thr

70

75

80

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90
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<211> 60
<212>
      DNA
<213>
      Mycobacterium tuberculosis
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<222>
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Met Ser Gln Ile Met Tyr Asn Tyr Pro Ala Met Met Ala His Ala Gly
                                    10
                                                                      60
gac atg gcc ggt
Asp Met Ala Gly
           20
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<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
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Met Ser Gln Ile Met Tyr Asn Tyr Pro Ala Met Met Ala His Ala Gly
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                5
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Asp Met Ala Gly
            20
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<221> CDS
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                                                                      48
atg atg get cat gee ggg gac atg gee ggt tat geg gge acg etg cag
Met Met Ala His Ala Gly Asp Met Ala Gly Tyr Ala Gly Thr Leu Gln
                                                                      60
agc ttg ggg gcc
Ser Leu Gly Ala
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Met Ala Met Leu Ala Arg Asp Gly Ala Glu Ala Ala Lys Trp Gly Gly

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<212> PRT
<213> Mycobacterium tuberculosis
<400> 203
Met Met Ala His Ala Gly Asp Met Ala Gly Tyr Ala Gly Thr Leu Gln
                                   10
Ser Leu Gly Ala
           20
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<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
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<221> CDS
<222>
      (1)..(60)
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tat gcg ggc acg ctg cag agc ttg ggg gcc gat atc gcc agt gag cag
Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Asp Ile Ala Ser Glu Gln
                                    10
                                                                      60
gcc gtg ctg tcc
Ala Val Leu Ser
           20
<210> 205
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 205
Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Asp Ile Ala Ser Glu Gln
Ala Val Leu Ser
            20
<210> 206
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
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<222> (1)..(60)

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gat atc gcc agt gag cag gcc gtg ctg tcc agt gct tgg cag ggt gat
                                                                     48
Asp Ile Ala Ser Glu Gln Ala Val Leu Ser Ser Ala Trp Gln Gly Asp
acc ggg atc acg
                                                                     60
Thr Gly Ile Thr
           20
<210> 207
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> -207
Asp Ile Ala Ser Glu Gln Ala Val Leu Ser Ser Ala Trp Gln Gly Asp
                                   10
Thr Gly Ile Thr
<210> 208
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222>
      (1)..(60)
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                                                                     48
agt gct tgg cag ggt gat acc ggg atc acg tat cag ggc tgg cag acc
Ser Ala Trp Gln Gly Asp Thr Gly Ile Thr Tyr Gln Gly Trp Gln Thr
                                   10
                                                                      60
cag tgg aac cag
Gln Trp Asn Gln
           20
<210> 209
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 209
Ser Ala Trp Gln Gly Asp Thr Gly Ile Thr Tyr Gln Gly Trp Gln Thr
1
               5
                             10
                                                      15
```

Gln Trp Asn Gln

L = n' = L

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<210> 210
<211> 60
<212> DNA
<213>
      Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(60)
<400> 210
tat cag ggc tgg cag acc cag tgg aac cag gcc cta gag gat ctg gtg
                                                                      48
Tyr Gln Gly Trp Gln Thr Gln Trp Asn Gln Ala Leu Glu Asp Leu Val
                                    10
                                                                       60
cgg gcc tat cag
Arg Ala Tyr Gln
            20
<210> 211
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 211
Tyr Gln Gly Trp Gln Thr Gln Trp Asn Gln Ala Leu Glu Asp Leu Val
                                    10
Arg Ala Tyr Gln
            20
<210> 212
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222>
      (1)..(60)
<400> 212
gcc cta gag gat ctg gtg cgg gcc tat cag tcg atg tct ggc acc cat
                                                                       48
Ala Leu Glu Asp Leu Val Arg Ala Tyr Gln Ser Met Ser Gly Thr His
                                    10
                                                         15
1
                5
                                                                       60
gag tcc aac acc
Glu Ser Asn Thr
            20
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<210> 213

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<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 213
Ala Leu Glu Asp Leu Val Arg Ala Tyr Gln Ser Met Ser Gly Thr His
               5
                                   10
Glu Ser Asn Thr
           20
<210> 214
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(60)
<400> 214
tcg atg tct ggc acc cat gag tcc aac acc atg gcg atg ttg gct cga
                                                                     48
Ser Met Ser Gly Thr His Glu Ser Asn Thr Met Ala Met Leu Ala Arg
                                   10
                                                                      60
gat ggg gcc gaa
Asp Gly Ala Glu
           20
<210> 215
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 215
Ser Met Ser Gly Thr His Glu Ser Asn Thr Met Ala Met Leu Ala Arg
                                   10
Asp Gly Ala Glu
<210> 216
<211> 48
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
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<222> (1)..(48)

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1 19 3
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<400> 216

<220>

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48
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Met Ala Met Leu Ala Arg Asp Gly Ala Glu Ala Ala Lys Trp Gly Gly
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<211> 16
<212> PRT
<213> Mycobacterium tuberculosis
<400> 217
Met Ala Met Leu Ala Arg Asp Gly Ala Glu Ala Ala Lys Trp Gly Gly
                                    10
                5
<210> 218
<211> 54
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(54)
<400> 218
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                                                                      48
Met Ser Gln Ile Met Tyr Asn Tyr Pro Ala Met Leu Gly His Ala Gly
                                    10
                                                        15
               5
                                                                      54
gat atg
Asp Met
<210> 219
<211> 18
<212> PRT
<213> Mycobacterium tuberculosis
<400> 219
Met Ser Gln Ile Met Tyr Asn Tyr Pro Ala Met Leu Gly His Ala Gly
                5
                                    10
                                                        15
Asp Met
<210> 220
<211> 54
<212> DNA
<213> Mycobacterium tuberculosis
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```
<221> CDS
<222> (1)..(54)
<400> 220
                                                                      48
atg ttg ggt cac gcc ggg gat atg gcc gga tat gcc ggc acg ctg cag
Met Leu Gly His Ala Gly Asp Met Ala Gly Tyr Ala Gly Thr Leu Gln
                                    10
                                                                      54
agc ttg
Ser Leu
<210> 221
<211> 18
<212> PRT
<213> Mycobacterium tuberculosis
<400> 221
Met Leu Gly His Ala Gly Asp Met Ala Gly Tyr Ala Gly Thr Leu Gln
Ser Leu
<210> 222
<211> 54
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
      (1)..(54)
<222>
<400> 222
                                                                      48
tat gcc ggc acg ctg cag agc ttg ggt gcc gag atc gcc gtg gag cag
Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Glu Ile Ala Val Glu Gln
                                                        15
                                    10
                                                                      54
gcc gcg
Ala Ala
<210> 223
<211> 18
<212> PRT
<213> Mycobacterium tuberculosis
<400> 223
Tyr Ala Gly Thr Leu Gln Ser Leu Gly Ala Glu Ile Ala Val Glu Gln
                5
                                    10
                                                        15
```

```
Ala Ala
```

```
<210> 224
<211> 54
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222>
      (1)..(54)
<400> 224
                                                                      48
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Glu Ile Ala Val Glu Gln Ala Ala Leu Gln Ser Ala Trp Gln Gly Asp
                                                                      54
acc ggg
Thr Gly
<210> 225
<211> 18
<212> PRT
<213> Mycobacterium tuberculosis
<400> 225
Glu Ile Ala Val Glu Gln Ala Ala Leu Gln Ser Ala Trp Gln Gly Asp
                                    10
Thr Gly
<210> 226
<211> 54
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221>
      CDS
<222>
      (1)..(54)
<400> 226
                                                                      48
agt gcg tgg cag ggc gat acc ggg atc acg tat cag gcg tgg cag gca
Ser Ala Trp Gln Gly Asp Thr Gly Ile Thr Tyr Gln Ala Trp Gln Ala
                                                                      54
cag tgg
Gln Trp
```

```
<210> 227
<211> 18
<212> PRT
<213> Mycobacterium tuberculosis
<400> 227
Ser Ala Trp Gln Gly Asp Thr Gly Ile Thr Tyr Gln Ala Trp Gln Ala
                                   10
Gln Trp
<210> 228
<211> 51
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(51)
<400> 228
                                                                     48
tat cag gcg tgg cag gca cag tgg aac cag gcc atg gaa gat ttg gtg
Tyr Gln Ala Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Asp Leu Val
                                                                      51
cgg
Arg
<210> 229
<211> 17
<212> PRT
<213> Mycobacterium tuberculosis
<400> 229
Tyr Gln Ala Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Asp Leu Val
                                    10
Arg
<210> 230
<211> 54
<212> DNA
<213> Mycobacterium tuberculosis
<220>
```

<221> CDS

```
<222> (1)..(54)
<400> 230
gcc atg gaa gat ttg gtg cgg gcc tat cat gcg atg tcc agc acc cat
                                                                      48
Ala Met Glu Asp Leu Val Arg Ala Tyr His Ala Met Ser Ser Thr His
               5
                                                                      54
gaa gcc
Glu Ala
<210> 231
<211> 18
<212> PRT
<213> Mycobacterium tuberculosis
<400> 231
Ala Met Glu Asp Leu Val Arg Ala Tyr His Ala Met Ser Ser Thr His
                                    10
Glu Ala
<210> 232
<211> 54
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(54)
<400> 232
gcg atg tcc agc acc cat gaa gcc aac acc atg gcg atg atg gcc cgc
                                                                      48
Ala Met Ser Ser Thr His Glu Ala Asn Thr Met Ala Met Met Ala Arg
                                                                      54
gac acg
Asp Thr
<210> 233
<211> 18
<212> PRT
<213> Mycobacterium tuberculosis
<400> 233
Ala Met Ser Ser Thr His Glu Ala Asn Thr Met Ala Met Met Ala Arg
                                    10
```

```
Asp Thr
```

```
<210> 234
<211> 48
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(48)
<400> 234
                                                                     48
atg gcg atg atg gcc cgc gac acc gcc gaa gcc gcc aaa tgg ggc ggc
Met Ala Met Met Ala Arg Asp Thr Ala Glu Ala Ala Lys Trp Gly Gly
                                    10
<210> 235
<211> 16
<212> PRT
<213> Mycobacterium tuberculosis
<400> 235
Met Ala Met Met Ala Arg Asp Thr Ala Glu Ala Ala Lys Trp Gly Gly
<210> 236
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(60)
<400> 236
                                                                      48
gtg tcg cag agt atg tac agc tac ccg gcg atg acg gcc aat gtc gga
Val Ser Gln Ser Met Tyr Ser Tyr Pro Ala Met Thr Ala Asn Val Gly
                                                        15
1
                                                                      60
gac atg gcc ggt
Asp Met Ala Gly
            20
<210> 237
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 237
Val Ser Gln Ser Met Tyr Ser Tyr Pro Ala Met Thr Ala Asn Val Gly
```

```
1 5 10 15
```

```
Asp Met Ala Gly
            20
<210> 238
<211> 60
<212>
      DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222>
      (1)..(60)
<400> 238
atg acg gcc aat gtc gga gac atg gcc ggt tat acg ggc acg acg cag
                                                                       48
Met Thr Ala Asn Val Gly Asp Met Ala Gly Tyr Thr Gly Thr Thr Gln
                                    10
                                                                       60
agc ttg ggg gcc
Ser Leu Gly Ala
            20
<210> 239
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 239
Met Thr Ala Asn Val Gly Asp Met Ala Gly Tyr Thr Gly Thr Thr Gln
Ser Leu Gly Ala
            20
<210> 240
<211> 60
<212> DNA
      Mycobacterium tuberculosis
<220>
<221>
      CDS
<222>
       (1)..(60)
<400> 240
tat acg ggc acg acg cag agc ttg ggg gcc gat atc gcc agt gag cgc
                                                                       48
Tyr Thr Gly Thr Thr Gln Ser Leu Gly Ala Asp Ile Ala Ser Glu Arg
                5
                                    10
                                                        15
acc gcg ccg tcg
                                                                       60
```

Thr Ala Pro Ser

```
<210> 241
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 241
Tyr Thr Gly Thr Thr Gln Ser Leu Gly Ala Asp Ile Ala Ser Glu Arg
Thr Ala Pro Ser
           20
<210> 242
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(60)
<400> 242
gat atc gcc agt gag cgc acc gcg ccg tcg cgt gct tgc caa ggt gat
                                                                     48
Asp Ile Ala Ser Glu Arg Thr Ala Pro Ser Arg Ala Cys Gln Gly Asp
                                                                      60
ctc ggg atg agt
Leu Gly Met Ser
           20
<210> 243
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 243
Asp Ile Ala Ser Glu Arg Thr Ala Pro Ser Arg Ala Cys Gln Gly Asp
                                    10
Leu Gly Met Ser
           20
<210> 244
<211> 60
<212>
      DNA
<213> Mycobacterium tuberculosis
```

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<220>
<221> CDS
<222> (1)..(60)
<400> 244
cgt gct tgc caa ggt gat ctc ggg atg agt cat cag gac tgg cag gcc
                                                                     48
Arg Ala Cys Gln Gly Asp Leu Gly Met Ser His Gln Asp Trp Gln Ala
               5
                                                                     60
cag tgg aat cag
Gln Trp Asn Gln
           20
<210> 245
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 245
Arg Ala Cys Gln Gly Asp Leu Gly Met Ser His Gln Asp Trp Gln Ala
      5
Gln Trp Asn Gln
<210> 246
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222> (1)..(60)
cat cag gac tgg cag gcc cag tgg aat cag gcc atg gag gct ctc gcg
                                                                     48
His Gln Asp Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Ala Leu Ala
                                                                     60
cgg gcc tac cgt
Arg Ala Tyr Arg
<210> 247
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 247
His Gln Asp Trp Gln Ala Gln Trp Asn Gln Ala Met Glu Ala Leu Ala
                                   10
```

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Arg Ala Tyr Arg
           20
<210> 248
<211>
      60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222>
      (1)..(60)
<400> 248
gcc atg gag gct ctc gcg cgg gcc tac cgt cgg tgc cgg cga gca cta
                                                                      48
Ala Met Glu Ala Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu
                                    10
                                                                      60
cgc cag atc ggg
Arg Gln Ile Gly
           20
<210> 249
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 249
Ala Met Glu Ala Leu Ala Arg Ala Tyr Arg Arg Cys Arg Arg Ala Leu
                                    10
Arg Gln Ile Gly
<210> 250
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222>
      (1)..(60)
<400> 250
cgg tgc cgg cga gca cta cgc cag atc ggg gtg ctg gaa agg ccg gta
                                                                      48
Arg Cys Arg Arg Ala Leu Arg Gln Ile Gly Val Leu Glu Arg Pro Val
                                    10
ggc gat tcg tca
                                                                      60
Gly Asp Ser Ser
```

```
<210> 251
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 251
Arg Cys Arg Arg Ala Leu Arg Gln Ile Gly Val Leu Glu Arg Pro Val
                                    10
Gly Asp Ser Ser
            20
<210> 252
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<220>
<221> CDS
<222>
      (1)..(60)
<400> 252
                                                                      48
gtg ctg gaa agg ccg gta ggc gat tcg tca gac tgc gga acg att agg
Val Leu Glu Arg Pro Val Gly Asp Ser Ser Asp Cys Gly Thr Ile Arg
                                    10
                                                                      60
gtg ggg tcg ttc
Val Gly Ser Phe
            20
<210> 253
<211> 20
<212> PRT
<213> Mycobacterium tuberculosis
<400> 253
Val Leu Glu Arg Pro Val Gly Asp Ser Ser Asp Cys Gly Thr Ile Arg
               5
                                    10
                                                        15
Val Gly Ser Phe
<210> 254
<211> 60
<212> DNA
<213> Mycobacterium tuberculosis
<400> 254
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60 gactgcggaa cgattagggt ggggtcgttc cggggtcggt ggctggaccc gcgccatgcg <210> 255 <211> 20 <212> PRT <213> Mycobacterium tuberculosis <400> 255 Asp Cys Gly Thr Ile Arg Val Gly Ser Phe Arg Gly Arg Trp Leu Asp Pro Arg His Ala 20 <210> 256 <211> 60 <212> DNA <213> Mycobacterium tuberculosis <400> 256 cggggtcggt ggctggaccc gcgccatgcg ggtccagcca cggccgccga cgccggagac 60 <210> 257 <211> 20 <212> PRT <213> Mycobacterium tuberculosis <400> 257 Arg Gly Arg Trp Leu Asp Pro Arg His Ala Gly Pro Ala Thr Ala Ala Asp Ala Gly Asp 20

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